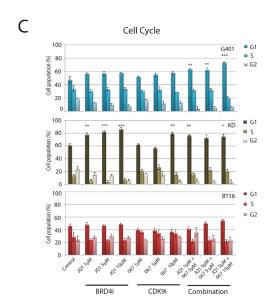
Combined BRD4 and CDK9 inhibition as a new therapeutic approach in malignant rhabdoid tumors

SUPPLEMENTARY FIGURES

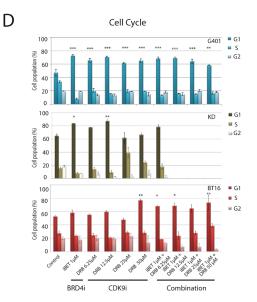
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Cell line	Treatment	IC50(μM)	R^2	CI
G401	JQ1 067 Combination	42.17 158.45 149.97	0.96 0.93 0.96	1.005
BT16	JQ1 067 Combination	23.43 39.28 7.83	0.99 0.99 0.97	0.217
KD	JQ1 067 Combination	42.67 142.6 103.83	0,97 0.99 0.97	1.005
A204	JQ1 067 Combination	30.36 52.19 23.84	0.93 0.95 0.93	0.771
MON	JQ1 067 Combination	40.98 92.52 23.64	0.99 0.94 0.93	0.56

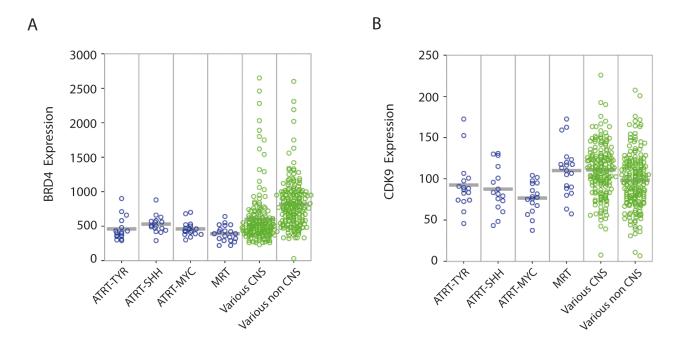


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Cell line	Treatment	IC50(μM)	R ²	CI
G401	iBET DRB Combination	23.99 376.99 8.95	0.84 0.71 0.82	0.05
BT16	iBET DRB Combination	189.4 65.21 56.64	0.98 0.99 0.91	0.373
KD	iBET DRB Combination	4.51 48.59 3.12	0.91 0.99 0.94	0.638
A204	iBET DRB Combination	27.07 142.85 10.6	0.99 0.97 0.93	0.365
MON	iBET DRB Combination	5.08 73.04 3.7	0.99 0.94 0.94	0.671



Supplementary Figure 1: Combined inhibition of BRD4 and CDK9 synergistically inhibits cell proliferation and induces cell cycle arrest in rhabdoid tumor cells in vitro. Different RT cell lines were incubated with BRD4 and CDK9 inhibitors at a range of concentrations $(0-100 \, \mu\text{M})$ as single compounds or as a combination with JQ1/ LDC067 (A) or iBET/DRB (B). IC₅₀ and CI values obtained by MTT assays are shown. Combined indexes (CI) were calculated using Chou-Thalalay's algorithm. CI indicates cooperation (synergistic or additive) between BRD4 and CDK9 inhibitors after simultaneous treatments. Cell cycle profile analyzed by flow cytometry after treatment with BRD4 and CDK9 inhibitors alone or together in the indicated cell lines is shown. Simultaneous application of JQ1/ LDC067 (C) or iBET/DRB (D) elicits a G1 Phase arrest. IC₅₀: Drug concentrations causing 50% growth inhibition, CI: Combination Index: indicates synergism if CI < 1, antagonism for CI > 1 and an additive effect for CI \approx 1. *p<0.05, **p < 0.01, ***p < 0.001 (ANOVA One-way Test).



Supplementary Figure 2: Expression of BRD4 and CDK9 in rhabdoid tumors. Gene expression profiling analysis of BRD4 **(A)** and CDK9 **(B)** performed on human samples of intracranial (AT/RT) and extracranial (MRT) tumors show a similar expression of both genes in tumors compared to healthy tissue.